

Claims

5 1. Method to distinguish, whether an event sequence is a memory driven event sequence or is not a memory driven event sequence on a time scale  $T_1$  to  $T_2$ , where  $T_1 < T_2$  are arbitrary times, characterized in that

10 a) the first order autocorrelation function  $G(\tau)$  of the event sequence is calculated,

b) the second order autocorrelation function  $G(\tau_1, \tau_2)$  of the event sequence is calculated,

15 c) it is decided that the event sequence is a memory driven event sequence on the time scale  $T_1$  to  $T_2$ ,

20 if the second order autocorrelation function of the event sequence can be expressed within experimental error as the product of first order autocorrelation functions, i.e.  $G(\tau_1, \tau_2) = G(\tau_1) * G(\tau_2)$  for  $T_1 < \tau_1, \tau_2 < T_2$ , and

25 d) it is decided that the event sequence is not a memory driven event sequence on the time scale  $T_1$  to  $T_2$ ,

if the second order autocorrelation function of the event sequence cannot be expressed within experimental error as the product of first order autocorrelation functions, i.e.  $G(\tau_1, \tau_2) \neq G(\tau_1) * G(\tau_2)$  for  $T_1 < \tau_1, \tau_2 < T_2$ .

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2. Method according to claim 1, characterized in that

- a) the first order autocorrelation function  $G(\tau)$  of the event sequence is calculated as:

$$G(\tau) \equiv \frac{E(X_0 X_\tau)}{E(X_0)E(X_\tau)}$$

where  $X$  is the random variable that describes the event and  $E(.)$  denotes the expectation value,

- b) the second order autocorrelation function  $G(\tau_1, \tau_2)$  of the event sequence is calculated as:

$$G(\tau_1, \tau_2) \equiv \frac{E(X_0 X_{\tau_1} X_{\tau_1+\tau_2})}{E(X_0)E(X_{\tau_1})E(X_{\tau_1+\tau_2})}$$

where  $X$  is the random variable that describes the event and  $E(.)$  denotes the expectation value,

3. Method according to claim 1, characterised in that the degree of memory of the system is quantified by the non-Markovian function NMF calculated according to:

$$\text{NMF}(\tau_1, \tau_2) = p_f \left( \frac{G(\tau_1, \tau_2)}{G(\tau_2)} - G(\tau_1) \right),$$

where  $p_f$  is the probability of the event  $X$  at a particular time.

4. Method according to claim 1, characterized in that the event sequence is a sequence of fluorescence events observed in a confocal microscope.

5. Method according to claim 4 to discriminate an event sequence from a single molecule against an event sequence from background processes or noise,

**characterized in that**

- a) it is decided that the event sequence is due to a single molecule, if it is a memory driven event sequence,
- b) it is decided that the event sequence is due to background processes or noise, if it is a non-memory driven event sequence.

6. Method according to claim 5 for single molecule sequencing,

**characterized in that**

- a) it is decided that the fluorescence events observed are due to nuclease-liberated nucleotides if the sequence of fluorescence events is a memory driven sequence of events and
- b) it is decided that the fluorescence events observed are due to contaminating nucleotides or other background signals, if the sequence of fluorescence events is not a memory driven sequence of events.

7. Method according to claim 6, characterized in that the fluorescence events are observed in a confocal microscope.

8. Method according to claim 6 or 7 for analyzing of catalytic complexes, characterized in that

- a) it is decided that the fluorescence events observed are due to characteristics of the catalytic complex if the sequence of fluorescence events is a memory driven sequence of events and
- b) it is decided that the fluorescence events observed are due to contaminating nucleotides or other background signals, if the

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sequence of fluorescence events is not a memory driven  
sequence of events.

5 9. Method according to claim 8, characterized in that the catalytic  
complex comprises a catalyst, a substrate being converted to a  
product and optionally a cosubstrate.

10 10. Method according to claim 8 or 9, characterized in that the catalyst  
is selected from biomolecules, e.g. enzymes, inorganic molecules  
and organic molecules.

11. Method according to one of the claims 5 - 10 wherein an oscillating  
process is analyzed.

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